

1/23

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MAY 1 2 2003

TECH CENTER 1600/2900

Gag AF110965 BW mod

ATGGGCGCCGCGCAGCATCCTGCGCGGCGCAAGCTGGACGCCTGGGAGCGCATCCGCC TGCGCCCGGCGGCAAGAAGTGCTACATGATGAAGCACCTGGTGTGGGCCAGCCGCGAGCT GGAGAAGTTCGCCCTGAACCCCGGCCTGCTGGAGACCAGCGAGGGCTGCAAGCAGATCATC CGCCAGCTGCACCCCGCCCTGCAGACCGGCAGCGAGGAGCTGAAGAGCCTGTTCAACACCG TGGCCACCCTGTACTGCGTGCACGAGAAGATCGAGGTCCGCGACACCAAGGAGGCCCTGGA CAAGATCGAGGAGGAGCAGAACAAGTGCCAGCAGAAGATCCAGCAGGCCGAGGCCGAC AAGGGCAAGGTGAGCCAGAACTACCCCATCGTGCAGAACCTGCAGGGCCAGATGGTGCACC AGGCCATCAGCCCCCGCACCCTGAACGCCTGGGTGAAGGTGATCGAGGAGAAGGCCTTCAG CCCCGAGGTGATCCCCATGTTCACCGCCCTGAGCGAGGGCGCCACCCCCCAGGACCTGAAC ACGATGTTGAACACCGTGGGCGGCCACCAGGCCGCCATGCAGATGCTGAAGGACACCATCA CCAGATGCGCGAGCCCCGCGGCAGCACCCGCACCACCAGCACCCTGCAGGAGCAG ATCGCCTGGATGACCAGCAACCCCCCCATCCCCGTGGGCGACATCTACAAGCGGTGGATCA TCCTGGGCCTGAACAAGATCGTGCGGATGTACAGCCCCGTGAGCATCCTGGACATCAAGCA GGGCCCAAGGAGCCCTTCCGCGACTACGTGGACCGCTTCTTCAAGACCCTGCGCGCCGAG CAGAGCACCCAGGAGGTGAAGAACTGGATGACCGACACCCTGCTGGTGCAGAACGCCAACC CCGACTGCAAGACCATCCTGCGCGCTCTCGGCCCCGGCGCCAGCCTGGAGGAGATGATGAC CGCCTGCCAGGCGTGGGCGGCCCCAGCCACAAGGCCCGCGTGCTGGCCGAGGCGATGAGC CAGGCCAACACCAGCGTGATGATGCAGAAGAGCAACTTCAAGGGCCCCCGGCGCATCGTCA AGTGCTTCAACTGCGGCAAGGAGGGCCACATCGCCCGCAACTGCCGCGCCCCCCGCAAGAA GGGCTGCTGGAAGTGCGGCAAGGAGGGCCACCAGATGAAGGACTGCACCGAGCGCCAGGCC AACTTCCTGGGCAAGATCTGGCCCAGCCACAAGGGCCGCCCCGGCAACTTCCTGCAGAGCC GCCCGAGCCCACCGCCCCCCCCGCGAGAGCTTCCGCTTCGAGGAGACCACCCCCGGCCA GAAGCAGGAGCAAGGACCGCGAGACCCTGACCAGCCTGAAGAGCCTGTTCGGCAACGAC CCCCTGAGCCAGTAA



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Gag_AF110967_BW mod ATGGGCGCCCGCGCAGCATCCTGCGCGCGAGAAGCTGGACAAGTGGGAGAAGATCCGCC TGCGCCCCGGCGCAAGAAGCACTACATGCTGAAGCACCTGGTGTGGGCCAGCCGCGAGCT GGAGGGCTTCGCCCTGAACCCCGGCCTGCTGGAGACCGCCGAGGGCTGCAAGCAGATCATG AAGCAGCTGCAGCCCGCCCTGCAGACCGGCAGCGAGGAGCTGCGCAGCCTGTACAACACCG TGGCCACCCTGTACTGCGTGCACGCCGGCATCGAGGTCCGCGACACCAAGGAGGCCCTGGA CAAGATCGAGGAGGAGCAGAACAAGTCCCAGCAGAAGACCCAGCAGGCCAAGGAGGCCGAC GGCAAGGTGAGCCAGAACTACCCCATCGTGCAGAACCTGCAGGGCCAGATGGTGCACCAGG CCATCAGCCCCGCACCCTGAACGCCTGGGTGAAGGTGATCGAGGAGAAGGCCTTCAGCCC CGAGGTGATCCCCATGTTCACCGCCCTGAGCGAGGGCGCCACCCCCAGGACCTGAACACG ATGTTGAACACCGTGGGCGGCCACCAGGCCGCCATGCAGATGCTGAAGGACACCATCAACG GATGCGCGACCCCCGCGGCACCACCAGCACCCTGCAGGAGCAGATC GCCTGGATGACCAGCAACCCCCCGTGCCCGTGGGCGACATCTACAAGCGGTGGATCATCC TGGGCCTGAACAAGATCGTGCGGATGTACAGCCCCGTGAGCATCCTGGACATCCGCCAGGG CCCCAAGGAGCCCTTCCGCGACTACGTGGACCGCTTCTTCAAGACCCTGCGCGCGAGCAG GCCACCCAGGACGTGAAGAACTGGATGACCGAGACCCTGCTGGTGCAGAACGCCAACCCCG ACTGCAAGACCATCCTGCGCGCTCTCGGCCCCCGGCGCCACCCTGGAGGAGATGATGACCGC $\tt CTGCCAGGGCGTGGGGGGGGGCCCACAAGGCCCGCGTGCTGGCCGAGGCGATGAGCCAG$ GCCAACAGCGTGAACATCATGATGCAGAAGAGCAACTTCAAGGGCCCCCGGCGCAACGTCA GGGCTGCTGGAAGTGCGGCAAGGAGGGCCACCAGATGAAGGACTGCACCGAGCGCCAGGCC AACTTCCTGGGCAAGATCTGGCCCAGCCACAAGGGCCGCCCCGGCAACTTCCTGCAGAACC GGAGACCACCCCCCCCAAGCAGGAGCCCAAGGACCGCGAGCCCTACCGCGAGCCCCTG ACCGCCCTGCGCAGCCTGTTCGGCAGCGGCCCCCTGAGCCAGTAA

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Polynucleotides Encoding Antigenic HIV Type C Polypeptides, Polypeptides and Uses Thereof

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Env_AF110968_C_BW_opt

--> signal peptide (1-81)
ATGCGCGTGATGGGCATCCTGGAAGAACTACCAGCAGTGGTGGATGTGGGGCATCCTGGGCTTCTGGATGCTGATCA \/--> gp120/140/160 (82)
TCAGCAGCGTGGTGGCAACCTGTGGGTGACCACCCT GACCCCAACCCCAGGAGATCGTGCTGGAGAACGTGACCGAGAACTTCAACATGTGGAAGAACGACATGGTGGACC AGATGCACGAGGACATCATCAGCCTGTGGGACCAGAGCCTGAAGCCCTGCGTGAAGCTGACCCCCTGTGCGTGAC CCTGAAGTGCCGCAACGTGAACGCCACCAACAACATCAACAGCATGATCGACAACAGCAACAAGGGCGAGATGAAG TGGTGCCCCTGCAGGGCAACAACAGCAACGAGTACCGCCTGATCAACTGCAACACCAGCGCCATCACCCAGGCCTG CAGACCTTCAACGGCACCGGCCCCTGCAACAACGTGAGCAGCGTGCAGTGCGCCCCACGGCATCAAGCCCGTGGTGA GCACCCAGCTGCTGAACGGCAGCCTGGCCAAGGGCGAGATCATCATCCGCAGCGAGAACCTGGCCAACAACGC GTGCGCATCGGCCCGGCCAGACCTTCTACGCCACCGGCGAGATCATCGGCGACATCCGCCAGGCCTACTGCATCA TCAACAAGACCGAGTGGAACAGCACCCTGCAGGGCGTGAGCAAGAAGCTGGAGGAGCACTTCAGCAAGAAGGCCAT TGCGACACCAGCCAGCTGTTCAACAGCACCTACAGCCCCAGCTTCAACGGCACCGAGAACAAGCTGAACGGCACCA TCACCATCACCTGCCGCATCAAGCAGATCATCAACATGTGGCAGAAGGTGGGCCGCGCCATGTACGCCCCCCCAT CGCCGGCAACCTGACCTGCGAGAGCAACATCACCGGCCTGCTGCTGACCCGCGACGGCGGCAAGACCGGCCCCAAC GACACCGAGATCTTCCGCCCGGCGGCGGCGACATGCGCGACAACTGGCGCAACGAGCTGTACAAGTACAAGGTGG TGGAGATCAAGCCCCTGGGCGTGGCCCCCACCGAGGCCAAGCGCCGCGTGGTGGAGCGCGAGAAGCGCGCGTGGG CATCGGCGCCGTGTTCCTGGGCTTCCTGGGCGCCCGCCAGCATCACCCTGACCGTG CAGGCCCGCCTGCTGAGCGGCATCGTGCAGCAGCAGCAACAACCTGCTGCGCGCCATCGAGGCCCAGCAGCACC TGCTGCAGCTGACCGTGTGGGGCATCAAGCAGCTGCAGACCCGCATCCTGGCCGTGGAGCGCTACCTGAAGGACCA GCAGCTGCTGGGCATCTGGGGCTGCAGCGGCAAGCTGATCTGCACCACCGCCGTGCCCTGGAACAGCAGCTGGAGC AACCGCAGCCACGACGAGATCTGGGACAACATGACCTGGATGCAGTGGGACCGCGAGATCAACAACTACACCGACA CCATCTACCGCCTGCTGGAGGAGAGCCAGAACCAGCAGGAGAAGAACGAGAAGGACCTGCTGGCCCTGGACAGCTG GCAGAACCTGTGGAACTGGTTCAGCATCACCAACTGGCTGTGGTACATCAAGATCTTCATCATGATCGTGGGCGGC CTGATCGGCCTGCGCATCATCTTCGCCGTGCTGAGCATCGTGAACCGCGTGCGCCAGGGCTACAGCCCCCTGCCCT CGGCCGCAGCATCCGCCTGGTGAGCGGCTTCCTGGCCCTGGCCTGGGACGACCTGCGCAGCCTGTTCAGC TACCACCGCCTGCGCGACTTCATCCTGATCGCCGCCCGCGTGCTGGAGCTGCTGGGCCAGCGCGGCTGGGAGGCCC TGAAGTACCTGGGCAGCCTGGTGCAGTACTGGGGCCTGGAGCTGAAGAAGAGCGCCATCAGCCTGCTGGACACCAT CGCCATCGCCGTGGCCGAGGGCACCGACCGCATCATCGAGTTCATCCAGCGCATCTGCCGCGCCATCCGCAACATC GP160, GP41 (2547) <--\
CCCCGCCGCATCCGCCAGGGCTTCGAGGCCGCCCTGCAGTAA

45.38 MAY 0 8 2003 THE PRODUM

03/010.313 Polynucleotides Encoding Antigenic HIV Type C Polypeptides. Polypeptides and Uses Thereof

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Env_AF110975_C_BW_opt --> signal peptide (1-72)
ATGCGCGTGCGCGGCATCCTGCGCAGCTGGCAGCAGCGGGCATCCTGGGCTTCTGGATCTGCAGCG 9p120/140/160 (72)
GCCTGGGCAACCTGTGACCGTGTACGACGGCGTGTGCCCGTGTGGCGCGAGGCCACCACCCTGTTCTGCGC CCCCAGGAGATCGAGCTGGACAACGTGACCGAGAACTTCAACATGTGGAAGAACGACATGGTGGACCAGATGCACG AGGACATCATCAGCCTGTGGGACCAGAGCCTGAAGCCCCGCGTGAAGCTGACCCCCCTGTGCGTGACCCTGAAGTG CACCAACTACAGCACCAACTACAGCAACACCATGAACGCCACCAGCTACAACAACAACACCACCGAGGAGATCAAG AACTGCACCTTCAACATGACCACCGAGCTGCGCGACAAGAAGCAGCAGGTGTACGCCCTGTTCTACAAGCTGGACA TCGTGCCCCTGAACAGCAACAGCGGGGTACCGCCTGATCAACTGCAACACCAGCGCCATCACCCAGGCCTGCCC CAAGGTGAGCTTCGACCCCATCCCATCCACTACTGCGCCCCGCCGGCTACGCCATCCTGAAGTGCAAGAACAAC ACCAGCAACGGCACCGGCCCCTGCCAGAACGTGAGCACCGTGCAGTGCACCCACGGCATCAAGCCCGTGGTGAGCA CTACACCATCATCGTGCACCTGAACGACAGCGTGGAGATCGTGTGCACCCGCCCAACAACAACACCCGCAAGGGC ATCCGCATCGGCCCGGCCAGACCTTCTACGCCACCGAGAACATCATCGGCGACATCCGCCAGGCCCACTGCAACA TCAGCGCCGGCGAGTGGAACAAGGCCGTGCAGCGCGTGAGCGCCAAGCTGCGCGAGCACTTCCCCAACAAGACCAT TGCAACACCAGCAAGCTGTTCAACAGCAGCTACAACGGCACCAGCTACCGCGGCACCGAGAGCAACAGCAGCATCA CGAGGGCAACATCACCTGCAGCAGCAGCATCACCGGCCTGCTGCTGGCCGGCGGCGGCCTGGACAACATCACC ACCGAGATCTTCCGCCCCAGGGCGGCGACATGAAGGACAACTGGCGCAACGAGCTGTACAAGTACAAGGTGGTGG AGATCAAGCCCCTGGGCGTGGCCCCCACCGAGGCCAAGCGCCGCGTGGTGGAGCGCGAGAAGCGCCGTGGGCAT CGGCGCCGTGATCTTCGGCTTCCTGGGCGCCGCCGGCAGCAACATGGGCGCCGCCAGCATCACCCTGACCGCCCAG TGCAGCTGACCGTGTGGGGCATCAAGCAGCTGCAGGCCCGCGTGCTGGCCATCGAGCGCTACCTGAAGGACCAGCA AAGACCCAGGGCGAGATCTGGGAGAACATGACCTGGATGCAGTGGGACAAGGAGATCAGCAACTACACCGGCATCA TCTACCGCCTGCTGGAGGAGGCCAGAACCAGCAGGAGCAGAACGAGAAGGACCTGCTGGCCCTGGACAGCCGCAA GP140(2022)<--\/
CAACCTGTGGAGCTGGTTCAACATCAGCAACTGGCTGGTACATCAAGATCTTCATCATGATCGTGGGCGGCCTG ${\tt ATCGGCCTGCGCATCATCTTCGCCGTGCTGAGCATCGTGAACCGCGTGCGCCAGGGCTACAGCCCCCTGAGCTTCC}$ CACCGCCTGCGCGACCTGATCCTGGTGACCGCCGCGTGGTGGAGCTGCTGGGCCGCAGCAGCCCCCGCGGCCTGC AGCGCGGCTGGGAGCCCTGAAGTACCTGGGCAGCCTGGTGCAGTACTGGGGCCTGGAGCTGAAGAAGAGCGCCAC gp160, gp41(2565)<--\
CGCGCCTTCTGCAACATCCCCCGCCGCGTGCGCCAGGGCTTCGAGGCCGCCCTGCAGTAA



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Gag_AF110965_BW_opt ATGGGCGCCGCGCGGCATCCTGCGCGGCGCAAGCTGGACGCCTGGGAGCGCATCCGCCTGCGCCCCGG CGGCAAGAAGTGCTACATGATGAAGCACCTGGTGTGGGCCAGCCGCGAGCTGGAGAAGTTCGCCCTGAACC ${\tt CCGGCCTGCTGGAGACCAGCTGCAGGCTGCAGCCGGCCTGCAGACCGGC}$ AGCGAGGAGCTGAAGAGCCTGTTCAACACCGTGGCCACCCTGTACTGCGTGCACGAGAAGATCGAGGTGCG CGACACCAAGGAGGCCCTGGACAAGATCGAGGAGGAGCAGAACAAGAGCCAGCAGAAGATCCAGCAGGCCG CAGGCCATCAGCCCCGCACCCTGAACGCCTGGGTGAAGGTGATCGAGGAGAAGGCCTTCAGCCCCGAGGT GATCCCCATGTTCACCGCCCTGAGCGAGGGCGCCACCCCCAGGACCTGAACACCATGCTGAACACCGTGG GCGGCCACCAGGCCGCCATGCAGATGCTGAAGGACACCATCAACGAGGAGGCCGCCGAGTGGGACCGCGTG CACCCGTGCACGCCGGCCCATCGCCCCGGGCCAGATGCGCGAGCCCCGGGCAGCGACATCGCCGGCAC CACCAGCACCCTGCAGGAGCAGATCGCCTGGATGACCAGCAACCCCCCATCCCGTGGGCGACATCTACA AGCGCTGGATCATCCTGGGCCTGAACAAGATCGTGCGCATGTACAGCCCCGTGAGCATCCTGGACATCAAG CCAGGAGGTGAAGAACTGGATGACCGACACCCTGCTGGTGCAGAACGCCAACCCCGACTGCAAGACCATCC TGCGCGCCTGGGCCCCGGCGCCAGCCTGGAGGAGATGATGACCGCCTGCCAGGGCGTGGGCGGCCCCAGC $\tt CACAAGGCCCGCGTGCTGGCCGAGGCCAGGCCAACACCAGCGTGATGATGCAGAAGAGCAACTT$ CAAGGGCCCCCGCCGCATCGTGAAGTGCTTCAACTGCGGCAAGGAGGGCCACATCGCCCGCAACTGCCGCG GCCAACTTCCTGGGCAAGATCTGGCCCAGCCACAAGGGCCGCCCCGGCAACTTCCTGCAGAGCCGCCCCGA GCCCACCGCCCCCCCGGCGAGAGCTTCCGCTTCGAGGAGACCACCCCCGGCCAGAAGCAGGAGAGCAAGG ACCGCGAGACCCTGACCAGCCTGAAGAGCCTGTTCGGCAACGACCCCCTGAGCCAGTAA

MAY 0 8 2000

Polynucleotides Encoding Antigenic HIV Type C Polypeptides, Polypeptides and Uses Thereof

NECFINED

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Gag_AF110967_BW_opt ATGGGCGCCGCGCCAGCATCCTGCGCGGGGAGAAGCTGGACAAGTGGGAGAAGATCCGCCTGCGCCCCGG CGGCAAGAAGCACTACATGCTGAAGCACCTGGTGTGGGGCCAGCCGCGAGCTGGAGGGCTTCGCCCTGAACC ACCGAGGAGCTGCGCAGCCTGTACAACACCGTGGCCACCCTGTACTGCGTGCACGCCGGCATCGAGGTGCG AGGAGGCCGACGGCAAGGTGAGCCAGAACTACCCCATCGTGCAGAACCTGCAGGGCCAGATGGTGCACCAG GCCATCAGCCCCGCACCCTGAACGCCTGGGTGAAGGTGATCGAGGAGAAGGCCTTCAGCCCCGAGGTGAT CCCCATGTTCACCGCCCTGAGCGAGGGCGCCACCCCCAGGACCTGAACACCATGGTGAACACCGTGGGCG CAGCACCCTGCAGGAGCAGATCGCCTGGATGACCAGCAACCCCCCGTGCCCGTGGGCGACATCTACAAGC qCrggatcatcctgggcctgaacaagatcgtgcqCatgtacagccccgtgagcatcctggacatccgccag GGACGTGAAGAACTGGATGACCGAGACCCTGCTGGTGCAGAACGCCAACCCCGACTGCAAGACCATCCTGC GCGCCCTGGGCCCCCCCGGGCGAGGAGATGATGACCGCCTGCCAGGGCGTGGGCGGCCCCGGCCAC AAGGCCCGCGTGCTGGCCGAGGCCATGAGCCAGCCCAACAGCGTGAACATCATGATGCAGAAGAGCAACTT CAAGGGCCCCCCCCCCAACGTGAAGTGCTTCAACTGCGGCAAGGAGGGCCACATCGCCAAGAACTGCCGCG CCCCCGCAAGAAGGCTGCTGGAAGTGCGGCAAGGAGGCCCACCAGATGAAGGACTGCACCGAGCGCCAG GCCAACTTCCTGGGCAAGATCTGGCCCAGCCACAAGGGCCGCCCGGCAACTTCCTGCAGAACCGCAGCGA CCAAGCAGGAGCCCAAGGACCGCGAGCCCTACCGCGAGCCCTGACCGCCCTGCGCAGCCTGTTCGGCAGC GGCCCCTGAGCCAGTAA



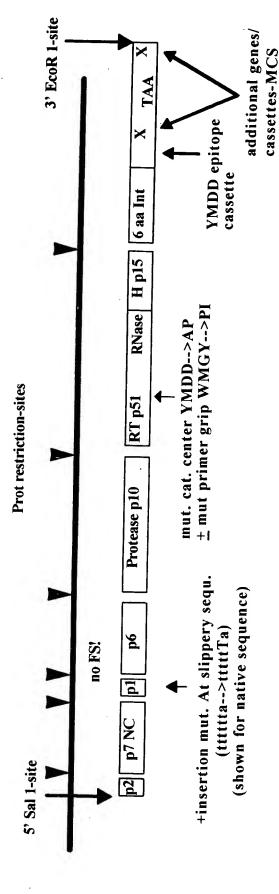


FIG. 7



8/23

PR975(+) (SEQ ID NO:30)

GTCGACGCCACCATGGCCGAGGCCATGAGCCAGGCCACCAGCGCCAACATCCTGAT GCAGCGCAGCAACTTCAAGGGCCCCAAGCGCATCATCAAGTGCTTCAACTGCGGCAA GGAGGGCCACATCGCCGCAACTGCCGCGCCCCCCGCAAGAAGGGCTGCTGGAAGT GCGGCAAGGAGGCCACCAGATGAAGGACTGCACCGAGCGCCAGGCCAACTTCTTC CGCGAGGACCTGGCCTTCCCCCAGGGCAAGGCCCGCGAGTTCCCCAGCGAGCAGAA CCGCGCCAACAGCCCACCAGCCGCGAGCTGCAGGTGCGCGGCGACAACCCCCGCA GCGAGGCCGGCCGAGCGCCAGGGCACCCTGAACTTCCCCCAGATCACCCTGTGGC AGCGCCCCTGGTGAGCATCAAGGTGGGCGGCCAGATCAAGGAGGCCCTGCTGGAC ACCGGCGCCGACGACACCGTGCTGGAGGAGATGAGCCTGCCCGGCAAGTGGAAGCC CAAGATGATCGGCGGCATCGGCGGCTTCATCAAGGTGCGCCAGTACGACCAGATCCT GATCGAGATCTGCGGCAAGAAGGCCATCGGCACCGTGCTGATCGGCCCCACCCCCGT GAACATCATCGGCCGCAACATGCTGACCCAGCTGGGCTGCACCCTGAACTTCCCCAT CAGCCCCATCGAGACCGTGCCCGTGAAGCTGAAGCCCGGCATGGACGGCCCCAAGG TGAAGCAGTGGCCCCTGACCGAGGAGAAGATCAAGGCCCTGACCGCCATCTGCGAG GAGATGGAGAAGGGGCAAGATCACCAAGATCGGCCCCGAGAACCCCTACAACAC CCCCGTGTTCGCCATCAAGAAGAAGGACAGCACCAAGTGGCGCAAGCTGGTGGACT TCCGCGAGCTGAACAAGCGCACCCAGGACTTCTGGGAGGTGCAGCTGGGCATCCCCC ACCCCGCCGGCCTGAAGAAGAAGAAGAGCGTGACCGTGCTGGACGTGGGCGACGCC TACTTCAGCGTGCCCCTGGACGAGGACTTCCGCAAGTACACCGCCTTCACCATCCCC AGCATCAACAACGAGACCCCCGGCATCCGCTACCAGTACAACGTGCTGCCCCAGGGC TGGAAGGCCAGCATCTTCCAGAGCAGCATGACCAAGATCCTGGAGCCCTTC CGCGCCCGCAACCCCGAGATCGTGATCTACCAGTACATGGACGACCTGTACGTGGGC AGCGACCTGGAGATCGGCCAGCACCGCGCCAAGATCGAGGAGCTGCGCAAGCACCT GCTGCGCTGGGGCTTCACCACCCCGACAAGAAGCACCAGAAGGAGCCCCCCTTCCT GTGGATGGGCTACGAGCTGCACCCCGACAAGTGGACCGTGCAGCCCATCGAGCTGCC CGAGAAGGAGAGCTGGACCGTGAACGACATCCAGAAGCTGGTGGGCAAGCTGAACT GGGCCAGCCAGATCTACCCCGGCATCAAGGTGCGCCAGCTGTGCAAGCTGCTGCGCG GCGCCAAGGCCCTGACCGACATCGTGCCCCTGACCGAGGAGGCCGAGCTGGAGCTG GCCGAGAACCGCGAGATCCTGCGCGAGCCCGTGCACGGCGTGTACTACGACCCCAG CAAGGACCTGGTGGCCGAGATCCAGAAGCAGGGCCACGACCAGTGGACCTACCAGA TCTACCAGGAGCCCTTCAAGAACCTGAAGACCGGCAAGTACGCCAAGATGCGCACC GCCCACACCAACGACGTGAAGCAGCTGACCGAGGCCGTGCAGAAGATCGCCATGGA GAGCATCGTGATCTGGGGCAAGACCCCCAAGTTCCGCCTGCCCATCCAGAAGGAGAC CTGGGAGACCTGGTGGACCGACTACTGGCAGGCCACCTGGATCCCCGAGTGGGAGTT CGTGAACACCCCCCCCCTGGTGAAGCTGTGGTACCAGCTGGAGAAGGAGCCCATCAT CGGCGCGAGACCTTCTACGTGGACGGCGCCCCAACCGCGAGACCAAGATCGGCA AGGCCGGCTACGTGACCGACCGGGGCCGGCAGAAGATCGTGAGCCTGACCGAGACC ACCAACCAGAAGACCGAGCTGCAGGCCATCCAGCTGGCCCTGCAGGACAGCGGCAG CGAGGTGAACATCGTGACCGACAGCCAGTACGCCCTGGGCATCATCCAGGCCCAGCC CGACAAGAGCGAGCGAGCTGGTGAACCAGATCATCGAGCAGCTGATCAAGAAGG AGAAGGTGTACCTGAGCTGGGTGCCCGCCCACAAGGGCATCGGCGGCAACGAGCAG ATCGACAAGCTGGTGAGCAAGGGCATCCGCAAGGTGCTGTTCCTGGACGGCATCGAT GGCGGCATCGTGATCTACCAGTACATGGACGACCTGTACGTGGGCAGCGGCGGCCCT AGGATCGATTAAAAGCTTCCCGGGGCTAGCACCGGTGAATTC



9/23

PR975YM (SEQ ID NO:31)

GTCGACGCCACCATGGCCGAGGCCATGAGCCAGGCCACCAGCGCCAACATCCTGAT GCAGCGCAGCAACTTCAAGGGCCCCAAGCGCATCATCAAGTGCTTCAACTGCGGCAA GGAGGCCACATCGCCCGCAACTGCCGCGCCCCCCGCAAGAAGGGCTGCTGGAAGT GCGGCAAGGAGGCCACCAGATGAAGGACTGCACCGAGCGCCAGGCCAACTTCTTC CGCGAGGACCTGGCCTTCCCCCAGGGCAAGGCCCGCGAGTTCCCCAGCGAGCAGAA CCGCGCCAACAGCCCACCAGCCGCGAGCTGCAGGTGCGCGGCGACAACCCCCGCA GCGAGGCCGCGCGAGCGCCAGGGCACCCTGAACTTCCCCCAGATCACCCTGTGGC AGCGCCCCTGGTGAGCATCAAGGTGGGCGGCCAGATCAAGGAGGCCCTGCTGGAC ACCGGCGCCGACGACACCGTGCTGGAGGAGATGAGCCTGCCCGGCAAGTGGAAGCC CAAGATGATCGGCGGCATCGGCGGCTTCATCAAGGTGCGCCAGTACGACCAGATCCT GATCGAGATCTGCGGCAAGAAGGCCATCGGCACCGTGCTGATCGGCCCCACCCCCGT GAACATCATCGGCCGCAACATGCTGACCCAGCTGGGCTGCACCCTGAACTTCCCCAT CAGCCCCATCGAGACCGTGCCCGTGAAGCTGAAGCCCGGCATGGACGGCCCCAAGG TGAAGCAGTGGCCCTGACCGAGGAGAAGATCAAGGCCCTGACCGCCATCTGCGAG GAGATGGAGAAGGGGCAAGATCACCAAGATCGGCCCCGAGAACCCCTACAACAC CCCCGTGTTCGCCATCAAGAAGAAGGACAGCACCAAGTGGCGCAAGCTGGTGGACT TCCGCGAGCTGAACAAGCGCACCCAGGACTTCTGGGAGGTGCAGCTGGGCATCCCCC ACCCCGCCGGCCTGAAGAAGAAGAAGAGCGTGACCGTGCTGGACGTGGGCGACGCC TACTTCAGCGTGCCCCTGGACGAGGACTTCCGCAAGTACACCGCCTTCACCATCCCC AGCATCAACAACGAGACCCCCGGCATCCGCTACCAGTACAACGTGCTGCCCCAGGGC TGGAAGGCAGCCCCAGCATCTTCCAGAGCAGCATGACCAAGATCCTGGAGCCCTTC CGCGCCCGCAACCCCGAGATCGTGATCTACCAGGCCCCCCTGTACGTGGGCAGCGAC CTGGAGATCGGCCAGCACCGCGCCAAGATCGAGGAGCTGCGCAAGCACCTGCTGCG CTGGGGCTTCACCACCCCGACAAGAAGCACCAGAAGGAGCCCCCCTTCCTGTGGAT GGGCTACGAGCTGCACCCCGACAAGTGGACCGTGCAGCCCATCGAGCTGCCCGAGA AGGAGAGCTGGACCGTGAACGACATCCAGAAGCTGGTGGGCCAAGCTGAACTGGGCC GAACCGCGAGATCCTGCGCGAGCCCGTGCACGGCGTGTACTACGACCCCAGCAAGG ACCTGGTGGCCGAGATCCAGAAGCAGGGCCACGACCAGTGGACCTACCAGATCTAC CAGGAGCCCTTCAAGAACCTGAAGACCGGCAAGTACGCCAAGATGCGCACCGCCCA CACCAACGACGTGAAGCAGCTGACCGAGGCCGTGCAGAAGATCGCCATGGAGAGCA TCGTGATCTGGGGCAAGACCCCCAAGTTCCGCCTGCCCATCCAGAAGGAGACCTGGG AGACCTGGTGGACCGACTACTGGCAGGCCACCTGGATCCCCGAGTGGGAGTTCGTGA ACACCCCCCCCTGGTGAAGCTGTGGTACCAGCTGGAGAAGGAGCCCATCATCGGCG CCGAGACCTTCTACGTGGACGCCCCCCCCAACCGCGAGACCAAGATCGGCAAGGCC GGCTACGTGACCGACCGGGCCGGCAGAAGATCGTGAGCCTGACCGAGACCACCAA TGAACATCGTGACCGACAGCCAGTACGCCCTGGGCATCATCCAGGCCCAGCCCGACA AGAGCGAGAGCGAGCTGGTGAACCAGATCATCGAGCAGCTGATCAAGAAGGAGAAG GTGTACCTGAGCTGGGCCCCCCCCCACAAGGGCATCGGCGCAACGAGCAGATCGA CAAGCTGGTGAGCAAGGGCATCCGCAAGGTGCTGTTCCTGGACGCCATCGATGGCG GCATCGTGATCTACCAGTACATGGACGACCTGTACGTGGGCAGCGGCGCCCTAGGA TCGATTAAAAGCTTCCCGGGGCTAGCACCGGTGAATTC



10/23

PR975YMWM (SEQ ID NO:32)

GTCGACGCCACCATGGCCGAGGCCATGAGCCAGGCCACCAGCGCCAACATCCTGAT GCAGCGCAGCAACTTCAAGGGCCCCAAGCGCATCATCAAGTGCTTCAACTGCGGCAA GGAGGGCCACATCGCCGCAACTGCCGCGCCCCCCGCAAGAAGGGCTGCTGGAAGT GCGGCAAGGAGGCCACCAGATGAAGGACTGCACCGAGCGCCAGGCCAACTTCTTC CGCGAGGACCTGGCCTTCCCCCAGGGCAAGGCCCGCGAGTTCCCCAGCGAGCAGAA CCGCGCCAACAGCCCACCAGCCGCGAGCTGCAGGTGCGCGGCGACAACCCCCGCA GCGAGGCCGGCGCGAGCGCCAGGGCACCCTGAACTTCCCCCAGATCACCCTGTGGC AGCGCCCCTGGTGAGCATCAAGGTGGGCGGCCAGATCAAGGAGGCCCTGCTGGAC ACCGGCGCCGACGACACCGTGCTGGAGGAGATGAGCCTGCCCGGCAAGTGGAAGCC CAAGATGATCGGCGGCATCGGCGGCTTCATCAAGGTGCGCCAGTACGACCAGATCCT GATCGAGATCTGCGGCAAGAAGGCCATCGGCACCGTGCTGATCGGCCCCACCCCCGT GAACATCATCGGCCGCAACATGCTGACCCAGCTGGGCTGCACCCTGAACTTCCCCAT CAGCCCCATCGAGACCGTGCCCGTGAAGCTGAAGCCCGGCATGGACGGCCCCAAGG TGAAGCAGTGGCCCCTGACCGAGGAGAAGATCAAGGCCCTGACCGCCATCTGCGAG GAGATGGAGAAGGGCCAAGATCACCAAGATCGGCCCCGAGAACCCCTACAACAC CCCCGTGTTCGCCATCAAGAAGAAGGACAGCACCAAGTGGCGCAAGCTGGTGGACT TCCGCGAGCTGAACAAGCGCACCCAGGACTTCTGGGAGGTGCAGCTGGGCATCCCCC ACCCCGCCGGCCTGAAGAAGAAGAAGAGCGTGACCGTGCTGGACGTGGGCGACGCC TACTTCAGCGTGCCCCTGGACGAGGACTTCCGCAAGTACACCGCCTTCACCATCCCC AGCATCAACAACGAGACCCCCGGCATCCGCTACCAGTACAACGTGCTGCCCCAGGGC TGGAAGGCCAGCATCTTCCAGAGCAGCATGACCAAGATCCTGGAGCCCTTC CGCGCCCGCAACCCCGAGATCGTGATCTACCAGGCCCCCCTGTACGTGGGCAGCGAC CTGGAGATCGGCCAGCACCGCGCCAAGATCGAGGAGCTGCGCAAGCACCTGCTGCG CTGGGGCTTCACCACCCCGACAAGAAGCACCAGAAGGAGCCCCCCTTCCTGCCCAT CGAGCTGCACCCCGACAAGTGGACCGTGCAGCCCATCGAGCTGCCCGAGAAGGAGA CGAGATCCTGCGCGAGCCCGTGCACGGCGTGTACTACGACCCCAGCAAGGACCTGGT GGCCGAGATCCAGAAGCAGGGCCACGACCAGTGGACCTACCAGATCTACCAGGAGC CCTTCAAGAACCTGAAGACCGGCAAGTACGCCAAGATGCGCACCCCACACCAAC GACGTGAAGCAGCTGACCGAGGCCGTGCAGAAGATCGCCATGGAGAGCATCGTGAT CTGGGGCAAGACCCCAAGTTCCGCCTGCCCATCCAGAAGGAGACCTGGGAGACCT GGTGGACCGACTACTGGCAGGCCACCTGGATCCCCGAGTGGGAGTTCGTGAACACCC CCCCCTGGTGAAGCTGTGGTACCAGCTGGAGAAGGAGCCCATCATCGGCGCCGAG ACCTTCTACGTGGACGGCGCCCAACCGCGAGACCAAGATCGGCAAGGCCGGCTA CGTGACCGACCGGGCCGGCAGAGATCGTGAGCCTGACCGAGACCACCAGA AGACCGAGCTGCAGGCCATCCAGCTGGCCCTGCAGGACAGCGGCAGCGAGGTGAAC ATCGTGACCGACAGCCAGTACGCCCTGGGCATCATCCAGGCCCAGCCCGACAAGAG CGAGAGCGAGCTGGTGAACCAGATCATCGAGCAGCTGATCAAGAAGGAGAAGGTGT ACCTGAGCTGGGTGCCCGCCCACAAGGGCATCGGCGGCAACGAGCAGATCGACAAG CTGGTGAGCAAGGCATCCGCAAGGTGCTGTTCCTGGACGGCATCGATGGCGGCATC GTGATCTACCAGTACATGGACGACCTGTACGTGGGCAGCGGCGGCCCTAGGATCGAT TAAAAGCTTCCCGGGGCTAGCACCGGTGAATTC



11/23

8_5_ZA (SEQ ID NO:33)

1 00/	3633 GGGTT					
1 10	GAAGGGTT A	ATTTACTCC A	AGAAAAGGC AI	AGAAATCCT TO	GATTTGTGG G	TCTATCACA
01 (CACAMOGCII (JITCCCTGAT (IGGCAAAACT 1	ACACACCGGG (CCAGGGGGG	
	TONCCITIOG	AIGGIGCTAC	AAGCTAGTGC	CAGTTGACCC	AGGGGACCTC	G3 3 G3 GG
	ACGGAGGAGA	AGACAACTGT	TTGCTACACC	CTATCACCCA	ACATCCA ACA	G3 GG
411	TINGAGAAGI	ATTAAAGTGG	AAGTTTGACA	GCCTCCTAGC	ACCCACACAC	3
J U I	AGCIACAICC	GGAGTATTAC	AAAGACTGCT	GACACAGAAG	CCACTTTCCC	COTTO
301	I CCMC I GGGG	CGTTCCGGGA	GGTGTGGTCT	GGGCGGGACT	TCCCACTCC	01100000
721	AIGCIGCAIA	TAAGCAGCTG	CTTTTCGCCT	GTACTGGGTC	TOTOTOTOTO	G1 GG1 G1
541	GAGCCTGGGA	GCCCTCTGGC	TATCTAGGGA	ACCCACTGCT	TAAGCCTCAA	TAAAGCTTGC
601	CTTGAGTGCT	TCTCCTACTC	TGTGCCCATC	TGTTGTGTGA	CTCTGGTAAC	TAGAGATCCC
661	TCAGACCCTT	ACACCACACA	TGGAAAATCT	CTAGCAGTGG	CGCCCGAACA	GGGACCAGAA
721	AGTGAAAGTG GGCAAGAGGC	GAGAGGGGGG	AGATCTCTCG	ACGCAGGACT	CGGCTTGCTG	AAGTGCACAC
781	GGCAAGAGGC GAAGGAGAGA	GATGGGTGCG	ACACCCTCA	ACGCCAATTT	TACTTGACTA	GCGGAGGCTA
841	GAAGGAGAGA AAAGAATTAG	GTTAAGGCCA	CCCCCAAAAA	TATTAAGCGG	CGGAAAATTA	GATAAATGGG
901	AAAGAATTAG CAAGCAGGGA	GCTGGAAAGA	TTTCCACTOR	AACATTATAT	GTTAAAACAT	CTAGTATGGG
961	CAAGCAGGGA GTAAACAAAT	AATAAAACAG	CTACAACCAA	ACCCTGGCCT	GTTAGAAACA	TCAGAAGGCT
1021	CATTATTCAA	САСАСТАССА	ACTOTOTA TO	CTCTTCAGAC	AGGAACAGAG	GAACTTAGAT
1081	CCAAGGAAGC	CTTAGACAAG	ACICICIAII	GIGTACATAA	AGGGATAGAG	GTACGAGACA
1141	AGGCAAAAGC	AGCTGACGAA	AIAGAGGAAG	AACAAAACAA	ATGTCAGCAA	AAAGCACAAC
1201	GGCAAATGGT	ACACCAAGCT	ARGGICAGIC	CAACATTATCC	TATAGTACAG	AATGCCCAAG
1261	AGGAAAAGGC	TTTCAATCCA	GAGGAAATAG	GAACATTGAA	TGCATGGATA	AAAGTAATAG
1321	CCCCACAAGA	TTTAAACACA	ATGTTAAATAC	CATGTTTAC	AGCATTATCA	GAAGGAGCCA
1381	TGTTAAAAGA	TACCATCAAT	GAGGAGGCTG	CAGIGGGGG	ACATCAAGCA	GCCATGCAAA
1441	CAGGGCCTGT	TGCACCAGGC	CAGATGAGAG	AACCAACCCC	TAGGACACAT	CCAGTACATG
1501	CTAGTACCCT	TCAGGAACAA	ATAGCATGGA	TGACAAGGGG	AAGTGACATA	GCAGGAACTA
1561	ACATCTATAA	AAGATGGATA	ATTCTGGGGT	TAACAAGIAA	TCCACCTATT	CCAGTAGAAG
1621	TTAGCATTTT	GGACATAAAA	CAAGGGCCAA	D D C D D C C C C C C C C C C C C C C C	AGTAAGAATG	TATAGCCCTG
1681	TCTTTAAAAC	CTTAAGAGCT	GAACAAGCTA	CACAACCCII	AAAGACTAT	GTAGACCGGT
1,741	CCTTGTTGGT	CCAAAATGCG	AACCCAGATT	GTAAGACCAT	AAAGAATIGG	ATGACAGACA
1001	GGGCCICAII	AGAAGAAATG	ATGACAGCAT	GTCAGGGAGT	CCCACCACCA	
7001	CAAGAGIGII	GGCTGAGGCA	ATGAGCCAAG	ሮልልልሮልሮሞአአ	CATACTACTAC	~~~~~
1721	VIIIIWWWGG	CICIAACAGA	ATTATTAAAT	ርጥጥጥር ል ልጣጥር	TCCCAAACONA	GGGG3
101	CCYGWWYIIG	CAGGGCCCCT	AGGAAAAAGG	GCTGTTGGAA	ATCTCCACAC	G3 3 GG3 G3 G3
2041	MANIGHAAGA	CIGIACIGAG	AGGCAGGCTA	אַ יייייייייייייייייייייייי	CAAAAMMMAA	aa
2101	AGGGGGGCC	AGGGAATTTC	CTCCAGAACA	GACCAGAGCC	A A C A C C C C C C A	aa
2101	CAACAGCCCC	ACCAGCAGAG	AGCTTCAGGT	TCGAGGAGAC	A A CCCCCCCCCC	0003.003.5.00
~ ~ ~ 1	DADADAGAGA	GGAACCTTTA	ACTTCCCTCA	<u>እ</u> እጥሮ እርጥር ውጥ	MODON COOK	
2201	WITWWWGIW	GAGGGCCAGA	TAAAGGAGGC	ጥርጥርጥጥአለ	3 C3 CC3 CC3 C	
2311	ATIAGAAGAA	ATAGATTTGC	CAGGGAAATG	GAAACCAAAA	ATCATACCCC	C3.3.000.000.00
	TITITICAMA	GIAAGACAGT	AIGAICAAAT	ልርጥጥልጥል ርአክ	እጥጥጣርመርፈጽጽ	
2401	MIDMINDO	TIAGTAGGGC	CTACACCAGT	ሮ ል ል ሮ ል ጥ ል ላ ጥጥ	CCARCARAMO	mamma =
2021	GCTIGGMIGC	ACACTAAATT	TTCCAATTAG	ע ע בטרויים ע ידייטיים א	A COMORNA CON C	
2301	ACCAGGAAIG	GAIGGCCCAA	AGGTCAAACA	Δ ጥር/ር/ር/ር/አ ጥጥ ር	707077077	
2011	VI IVVCHOCK	ATTIGICACC	AAATGGAGAA	CCAACCAAAA	3 MM3 43 5 5 5 5	
-, -,	TWICCHIMI	MACACICCAG	TATTTCCCAT	אמי אמת ממממ	03 03 0ms oms	
~,01	WI TWO I WOW!	LICAGGGAAC	TCAATAAAAA	<u>እ</u> ስርጥሮአ አረአረ	mmmaaa	
	MAINCONCAC	CCAGCAGGAT	עמבועם מעם מעם איוי	አአአአጥጣአጣጠጣ	3 63 65 65 65 6	
2881	TGCATATTTT	TCAGTTCCTT	TAGATGAAAG	CTTCAGGAAA	TATACTGCAT	TCACCATACC

FIG. 11A



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2941	TAGTATAAAC	AATGAAACAC	CAGGGATTAG	ATATCAATAT	AATGTGCTGC	CACAGGGATG
3001	GWWWGGWICW	CCAGCAATAT	TCCAGAGTAG	CATGACAAAA	ስጥርጥጥአ ለአ ለረ	
3001	WWWWWICCW	GACATAGTTA	TCTATCAATA	. TATGGATGA	שמויים מיתים מיתים ביית	G1.55555
3121	AGNAMIAGGG	CAACATAGAG	CAAAAATAGA	AGAGTTAAGG	CA A CA TOTO A TO	
2101	ATTIACAACA	CCAGACAAGA	. AACATCAAAA	AGAACCCCCA	עים ביים היים ביים א	TOCOCON
7271	MCICCAICCI	GACAAATGGA	CAGTACAACC	TATACTGCTG	CCAGAAAACC	A TO A COMPAGN -
3301	IGICAAIGAT	ATACAGAAGT	'TAGTGGGAAA	. ልጥጥልልል ርጥርር	GCDDCTCDCD	mmn
320T	GATTAAAGTA	AGGCAACTCT	GTAAACTCCT	'CAGGGGGGCC	AAAGCACTAA	CACACAMAG
3421	ACCACTAACT	GAAGAAGCAG	AATTAGAATT	GGCAGAGAAC	ልርርርል ል ል ጥጥ ጥ	TAACAGAAAG
240T	AGIACATGGA	GTATATTATG	ATCCATCAAA	ልርልርጥጥርልጥ ል	CCTCDDDDDDD	1011100000
334T	GCATGAACAA	TGGACATATC	AAATTTATCA	AGAACCATTT	AAAAATCTCA	AAACACCCA
360I	GIAIGCAAAA	ATGAGGACTA	CCCACACTAA	TGATGTAAAA	CAGTTAACAG	ACCCACMOON
2001	AAAAATAGCC	ATGGAAAGCA	TAGTAATATG	GGGAAAGACT	מידידים במבריים	GACTACCAN
3/21	CCAAAAAGAA	ACATGGGAGA	CATGGTGGAC	AGACTATTGG	CAAGCCACCT	GGATCGGTGA
3781	GTGGGAGTTT	GTTAATACCC	CTCCCCTAGT	AAAATTATGG	TACCAACTAG	AAAAAAA
3041	CATAGCAGGA	GTAGAAACTT	TCTATGTAGA	TGGAGCAACT	AATAGGGAAG	CTABBARAGA
3301	AAAAGCAGGG	TATGTTACTG	ACAGAGGAAG	GCAGAAAATT	GTTACTCTAA	CTARCACAR
3961	AAATCAGAAG	ACTGAGTTAC	AAGCAATTCA	GCTAGCTCTG	CAGGATTCAG	CAMCACAAC
4021	AAACATAGTA	ACAGACTCAC	AGTATGCATT	AGGAATCATT	CAAGCACAAC	CACAMAAGAG
4001	TGACTCAGAG	ATATTTAACC	AAATAATAGA	ACAGTTAATA	AACAAGGAAA	CA AUCUS COM
4141	GICAIGGGIA	CCAGCACATA	AAGGAATTGG	GGGAAATGAA	CAAGTAGATA	A A COURT A COURT A CO
4201	TAAGGGAATT	AGGAAAGTGT	TGTTTCTAGA	TGGAATAGAT	ADAGCTCAAC	AATTAGTAAG
4201	MAGGIACCAC	AGCAATTGGA	GAGCAATGGC	ጥ እጥር እር ጥጥጥ	እእጥርጥርርርርአር	CC3 E3 CE3 CE
422T	MAMAGAMATA	GTAGCTAGCT	GTGATAAATG	TCAGCTAAAA	GGGGAAGCCA	TACAMOOR OF
430T	AGICGACIGI	AGTCCAGGGA	TATGGCAATT	AGATTGTACC	CATTTAGAGG	CAAAAAMGAM
334T	CCIGGIAGCA	GICCATGTAG	CTAGTGGCTA	CATGGAAGCA	GAGGTTATCC	CACCACAAAA
#20T	AGGACAAGAA	ACAGCATATT	TTATATTAAA	ATTAGCAGGA	AGATGGCCAG	TCAAACTAAT
420T	ACATACAGAC	AATGGCAGTA	ATTTTACCAG	TACTGCAGTT	AAGGCAGCCT	GTTCCTCCCC
4021	AGGTATCCAA	CAGGAATTTG	GAATTCCCTA	CAATCCCCAA	AGTCAGGGAG	TOOTAGAARG
400T	CATGAATAAA	GAATTAAAGA	AAATAATAGG	ACAAGTAAGA	GATCAAGCTG	ACCA COMMA
4/41	GACAGCAGTA	CAAATGGCAG	TATTCATTCA	CAATTTTAAA	AGAAAAGGGG	CAATITICAGA
# 9 O T	GTACAGTGCA	GGGGAAAGAA	TAATAGACAT	AATAGCAACA	GACATACAAA	CM3 3 3 C3 3
4001	ACAAAAACAA	ATTATAAGAA	TTCAAAATTT	TCGGGTTTAT	TACAGAGACA	CCACACACAC
4721	INITIGGAAA	GGACCAGCCG	AACTACTCTG	GAAAGGTGAA	GGGGTAGTAG	TABTACTACT
4981	TAAAGGTGAC	ATAAAGGTAG	TACCAAGGAG	GAAAGCAAAA	ATCATTAGAG	ለ ጥጥ እ መረገር እ እ እ
3041	ACAGATGGCA	GGTGCTGATT	GTGTGGCAGG	TGGACAGGAT	GAAGATTAGA	CCATCCAAMA
2101	GITTAGTAAA	GCACCATATG	TATATATCAA	GGAGAGCTAG	TGGATGGGTC	TACACACATO
DIOI	ATTTTGAAAG	CAGACATCCA	AAAGTAAGTT	でなびななのでなべる	TATOOCA DON	~~~~~
5221	GATTAGTAAT	AAAAACATAT	TGGGGTTTGC	AGACAGGAGA	AAGAGATTCC	CATTOTOTOTO
T	MIGGWGICIC	CATAGAATGG	AGACTGAGAG	AATACAGCAC	ACA ACTACAC	CCTCAACCTCC
2241	CAGACCAGCT	AATTCACATG	CATTATTTTG	ልጥጥርጥጥጥል C	እርአ አጥርጥርርር	3003303030
240T	CCATALTAGG	ACACATAGTT	TTTCCTAGGT	GTGACTATCA	AGCAGGACAT.	AAGAAGGGAG
7401	GAICICIGCA	ATACTTGGCA	CTGACAGCAT	TGATAAAACC	AAAAAAAAAA	AACCCACCMC
JJZ1	IGCCIAGIGI	TAGAAAATTA	GTAGAGGATA	GATGGAACGA	ררררר ממאמ	ACCACCCCC
2201	GCAGAGGGAA	CCATACAATG	AATGGACACT	AGAGATTCTA	GDAGDACTCA	ACCACCA ACC
2047	TGTCAGACAC	TTTCCTAGAC	CATGGCTCCA	TAGCTTAGGA	ር እ አጥ አጥ አጥር ጥ	3 ma 3 3 a a a a
3701	IGGGGATACT	TGGACGGGAG	TTGAAGCTAT	AATAAGAGTA	でではて ひ か で か か ぐ	ጥ እ ርጥር መመረገ እ መ
3701	ICATITCAGA	ATTGGATGCC	AACATAGCAG	AATAGGCATC	TTCCCACACA	77777777
JU21	AAA1GGAGCC	AGTAGATCCT	AAACTAAAGC	CCTGGAACCA	TCCAGGAACC	C
5881	CAGCTTGTAA	TAATTGCTTT	TGCAAACACT	GTAGCTATCA	TTGTCTAGTT	ТССТОТОВСЬ
						COLLICAGA

FIG. 11B



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	•			•		
5941	CAAAAGGTTT	AGGCATTTCC	TATGGCAGGA	AGAAGCGGAG	ACAGCGACGA	AGCGCTCCTC
6001	CAAGTGGTGA	AGATCATCAA	AATCCTCTAT	CAAAGCAGTA	AGTACACATA	GTACATIONA A
6061	TGGTAAGTTT	AAGTTTATTT	AAAGGAGTAG	ATTATAGATT	AGGAGTAGGA	GC A TTC A TTA C
6121	TAGCACTAAT	CATAGCAATA	ATAGTGTGGA	CCATAGCATA	TATAGAATAT	ACC N N N TTTCC
6181	TAAGACAAAA	GAAAATAGAC	TGGTTAATTA	AAAGAATTAG	GGAAAGAGCA	GAAGAGAGTG
6241	GCAATGAGAG	TGATGGGGAC	ACAGAAGAAT	TGTCAACAAT	GGTGGATATC	CCCCATCTTA
6301	GGCTTCTGGA	TGCTAATGAT	TTGTAACACG	GAGGACTTGT	GGGTCACAGT	CTACTATGGG
6361	GTACCTGTGT	GGAGAGAAGC	AAAAACTACT	CTATTCTGTG	CATCAGATGG	TAAAGCATAT
6421	GAGACAGAAG	TGCATAATGT	CTGGGCTACA	CATGCTTGTG	TACCCACAGA	CCCCAACCCA
6481	CAAGAAATAG	TTTTGGGAAA	TGTAACAGAA	AATTTTAATA	TGTGGAAAA	TAACATGGCA
6541	GATCAGATGC	ATGAGGATAT	AATCAGTTTA	TGGGATCAAA	GCCTAAACCC	AMCMCMARA
6601	TTGACCCCAC	TCTGTGTCAC	TTTAAACTGT	ACAGATACAA	ATCTTACACC	MIGIGIAAAG
6661	GTTACAGGTA	ATACAAATGA	TACCAATATT	GCAAATGCTA	CATATATA	TAATAGAACT
6721	AAAAATTGCT	CTTTCAATGC	AACCACAGAA	TTAAGAGATA	CATATAAGTA	1GAAGAAATG
6781	CTCTTTTATA	AACTTGATAT	AGTACCACTT	DATGADATA	CTA A CA A COM	AGAGTATGCA
6841	TTAATAAATT	GCAATACCTC	AACCATAACA	CARCCCTCTC	GIAACAACTT	TACATATAGA
6901	ATTCCTATAC	ATTACTGTGC	TCCAGCTGAT	TATCCCATTC	CAAAGGTCTC	TTTTGACCCG
6961	TTCAATGGGA	CAGGACCATG	TTATAATCTC	ACCACACTAC	TAAAGTGTAA	TAATAAGACA
7021	CCAGTGGTAT	CAACTCAACT	Y CACAMAN Y WA	AGCACAGIAC	AATGTACACA	TGGAATTAAG
7081	AGATCTGAAA	ATTTGACAGA	GARRACCAAA	ACAAMAAMA	CAGAAGAAGG	GATAATAATT
7141	GAGATTAATT	GTACAAGGCC	CANTACCAAA	ACAATAATAG	TACATCTTAA	TGAATCTGTA
7201	CAAGCATTCT	ATGCAACAAA	TCACCARIAGI	ACAAGGAAAA	GTGTAAGGAT	AGGACCAGGA
7261	AGTACAGATA	GATGGAATAA	AACTTTTACAA	GGAAACATAA	GACAAGCACA	TTGTAACATT
7321	ССТААТАААА	CAIGGAAIAA	TCAACCACAC	CAGGTAATGA	AAAAATTAGG	AGAGCATTTC
7381	AGCTTTAATT	GTAGAGGAGA	ATTTTTTTTT	GCAGGAGGGG	ATCTAGAAAT	TACAATGCAT
7441	AGCTTTAATT TACTACCCTA	AGAATGGTAC	ATTITITITAT	1GCAATACAT	CAAACCTGTT	TAATAGTACA
7501	TACTACCCTA	TORATOGIAC	TCTACCCAMC	AATGGTAATT	CAAGCTTACC	CATCACACTC
7561	CAATGCAAAA	CACCAAACAT	AACATCTACA	TGGCAAGGGG	TAGGACAAGC	AATGTATGCC
7621	CCTCCCATTG	TTAACAACAC	AACAIGIAGA	TCAAACATCA	CAGGAATACT	ATTGACACGT
7681	GATGGGGGAT	ATAACTGCAC	A A CTICA A TITA	ACAGAGGAGA	CATTCAGACC	TGGAGGAGGA
7741	GATATGAGGG TTGGGAATAG	CACCCACTAA	CCCAAAAAA	TATAAATATA	AAGTGGTAGA	AATTAAGCCA
7801	TTGGGAATAG	CTCTCTTCCT	TCCCTTCTTC	AGAGTGGTGC	AGAGAAAAA	AAGAGCAGTG
7861	GGAATAGGAG TCAATAACGC	TGACGGTACA	CCCCACACA	GGAGCAGCAG	GAAGCACTAT	GGGCGCAGCG
7921	TCAATAACGC	ACCCTATACA	GGCCAGACAA	CTGTTGTCTG	GTATAGTGCA	ACAGCAAAGC
7981	AATTTGCTGA	ACCCCACACA	GGCGCAACAG	CATATGTTGC	AACTCACAGT	CTGGGGCATT
8041	AAGCAGCTCC	AGGCGAGAGT	CCTGGCTATA	GAAAGATACC	TAAAGGATCA	ACAGCTCCTA
8101	GGGATTTGGG	AAMGMGAAGG	AAGACTCATC	TGCACCACTG	CTGTGCCTTG	GAACTCCAGT
8161	TGGAGTAATA	AATCIGAAGC	AGATATTTGG	GATAACATGA	CTTGGATGCA	GTGGGATAGA
8221	GAAATTAATA	ATTACACAGA	AACAATATTC	AGGTTGCTTG	AAGACTCGCA	AAACCAGCAG
0221	GAAAAGAATG	AAAAAGATTT	ATTAGAATTG	GACAAGTGGA	ATAATCTGTG	GAATTGGTTT
0201	GACATATCAA	ACTGGCTGTG	GTATATAAAA	ATATTCATAA	TGATAGTAGG	AGGCTTGATA
0.401	GGTTTAAGAA	TAATTTTTGC	TGTGCTCTCT	ATAGTGAATA	GAGTTAGGCA	GGGATACTCA
0401	CCTTTGTCAT	TTCAGACCCT	TACCCCAAGC	CCGAGGGGAC	TCGACAGGCT	CGGAGGAATC
0401	GAAGAAGAAG	GTGGAGAGCA	AGACAGAGAC	AGATCCATAC	GATTGGTGAG	CGGATTCTTG
027T	TCGCTTGCCT	GGGACGATCT	GCGGAGCCTG	TGCCTCTTCA	GCTACCACCG	CTTGAGAGAC
8281	TTCATATTAA	TTGCAGTGAG	GGCAGTGGAA	CTTCTGGGAC	ACAGCAGTCT	CACCCCACTA
8641	CAGAGGGGGT	GGGAGATCCT	TAAGTATCTG	GGAAGTCTTG	TGCAGTATTG	GGGTCTAGAG
8,01	CTAAAAAAAGA	GTGCTATTAG	TCCGCTTGAT	ACCATAGCAA	TAGCAGTAGC	TGAAGGAACA
8/61	GATAGGATTA	TAGAATTGGT	ACAAAGAATT	TGTAGAGCTA	TCCTCAACAT	ACCTAGGAGA
8821	ATAAGACAGG	GCTTTGAAGC	AGCTTTGCTA	TAAAATGGGA	GGCAAGTGGT	CDADACCCAC
8881	CATAGTTGGA	TGGCCTGCAG	TAAGAGAAAG	AATGAGAAGA	ACTGAGCCAG	CAGCAGAGGG
8941	AGTAGGAGCA	GCGTCTCAAG	ACTTAGATAG	ACATGGGGCA	CTTACAAGCA	GCAACACACC



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9001	TGCTACTAAT	GAAGCTTGTG	CCTGGCTGCA	AGCACAAGAG	GAGGACGGAG	ATGTAGGCTT
9061	TCCAGTCAGA	CCTCAGGTAC	CTTTAAGACC	AATGACTTAT	AAGAGTGCAG	TAGATCTCAG
9121	CTTCTTTTTA	AAAGAAAAGG	GGGGACTGGA	AGGGTTAATT	TACTCTAGGA	AAAGGCAAGA
9181	AATCCTTGAT	TTGTGGGTCT	ATAACACACA	AGGCTTCTTC	CCTGATTGGC	AAAACTACAC
9241	ATCGGGGCCA	GGGGTCCGAT	TCCCACTGAC	CTTTGGATGG	TGCTTCAAGC	TAGTACCAGT
9301	TGACCCAAGG	GAGGTGAAAG	AGGCCAATGA	AGGAGAAGAC	AACTGTTTGC	TACACCCTAT
9361	GAGCCAACAT	GGAGCAGAGG	ATGAAGATAG	AGAAGTATTA	AAGTGGAAGT	TTGACAGCCT
9421	TCTAGCACAC	AGACACATGG	CCCGCGAGCT	ACATCCGGAG	TATTACAAAG	ACTGCTGACA
9481	CAGAAGGGAC	TTTCCGCCTG	GGACTTTCCA	CTGGGGCGTT	CCGGGAGGTG	TGGTCTGGGC
9541	GGGACTTGGG	AGTGGTCACC	CTCAGATGCT	GCATATAAGC	AGCTGCTTTT	CGCTTGTACT
9601	GGGTCTCTCT	CGGTAGACCA	GATCTGAGCC	TGGGAGCTCT	CTGGCTATCT	AGGGAACCCA
9661	CTGCTTAGGC	CTCAATAAAG	CTTGCCTTGA	GTGCTCTAAG	TAGTGTGTGC	CCATCTGTTG
9721	TGTGACTCTG	GTAACTAGAG	ATCCCTCAGA	CCCTTTGTGG	TAGTGTGGAA	AATCTCTAGC
0701	7.					CIAGC

FIG. 11D



15/23

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TECH CENTER 1600/2900

SEQ ID NO:34



16/23

975Pol wt until 6aa Int: (SEQ ID NO:35)

TTTTTTAGGGAAGATTTGGCCTTCCCACAAGGGAAGGCCAGGGAATTTCCTTCAGAA CAGAACAGACCCACCACCAGCAGAGAGCTTCAAGTTCGAGGAGACAACCC CCGCTCCGAAGCAGGAGCCGAAAGACAGGGAACCCTTAATTTCCCTCAAATCACTCT TTGGCAGCGACCCCTTGTCTCAATAAAAGTAGGGGGTCAAATAAAGGAGGCTCTCTT AGACACAGGAGCTGATGATACAGTATTAGAAGAAATGAGTTTGCCAGGAAAATGGA AACCAAAAATGATAGGAGGAATTGGAGGTTTTATCAAAGTAAGACAGTATGATCAA ATACTTATAGAAATTTGTGGAAAAAAGGCTATAGGTACAGTATTAATAGGACCTACA CCTGTCAACATAATTGGAAGGAATATGTTGACTCAGCTTGGATGCACACTAAATTTT AAGGTTAAACAATGGCCATTGACAGAAGAGAAAATAAAAGCATTAACAGCAATTTG TGAAGAAATGGAGAAAGGAAAAATTACAAAAATTGGGCCTGAAAATCCATATA ACACTCCAGTATTTGCCATAAAAAAGAAGGACAGTACTAAGTGGAGAAAGTTAGTA GATTTCAGGGAACTTAATAAAAGAACTCAAGACTTTTGGGAAGTTCAATTAGGAATA CCACACCCAGCAGGGTTAAAAAAAGAAAAAATCAGTGACAGTACTGGATGTGGGGGA TGCATATTTTTCAGTTCCTTTAGATGAGGACTTCAGGAAATATACTGCATTCACCATA CCTAGTATAAACAATGAAACACCAGGGATTAGATATCAATATAATGTGCTTCCACAG GGATGGAAAGGATCACCATCAATATTCCAGAGTAGCATGACAAAAATCTTAGAGCC CTTTAGAGCAAGAAATCCAGAAATAGTCATCTATCAATATATGGATGACTTGTATGT AGGATCTGACTTAGAAATAGGGCAACATAGAGCAAAAATAGAGGAGTTAAGAAAAC TTTCTTTGGATGGGTATGAACTCCATCCTGACAAATGGACAGTACAGCCTATAGAG TTGCCAGAAAAGGAAAGCTGGACTGTCAATGATATACAGAAGTTAGTGGGAAAATT AAATTGGGCCAGTCAGATTTACCCAGGAATTAAAGTAAGGCAACTTTGTAAACTCCT TAGGGGGCCAAAGCACTAACAGATATAGTACCACTAACTGAAGAAGCAGAATTAG AATTGGCAGAGAACAGGGAAATTCTAAGAGAACCAGTACATGGAGTATATTATGAC CCATCAAAAGACTTGGTAGCTGAAATACAGAAACAGGGGCATGACCAATGGACATA TCAAATTTACCAAGAACCATTCAAAAACCTGAAAACAGGGAAGTATGCAAAAATGA GGACTGCCCACACTAATGATGTAAAACAGTTAACAGAGGCAGTGCAAAAAATAGCT ATGGAAAGCATAGTAATATGGGGAAAGACTCCTAAATTTAGACTACCCATCCAAAA AGAAACATGGGAGACATGGTGGACAGACTATTGGCAAGCCACCTGGATTCCTGAGT CCATAATAGGAGCAGAAACTTTCTATGTAGATGGAGCAGCTAATAGGGAAACTAAA ATAGGAAAAGCAGGGTATGTTACTGACAGAGGAAGGCAGAAAATTGTTTCTCTAAC AGGATCAGAAGTAAACATAGTAACAGACTCACAGTATGCATTAGGAATCATTCAAG CACAACCAGATAAGAGTGAATCAGAGTTAGTCAACCAAATAATAGAACAATTAATA AAAAAGGAAAAGGTCTACCTGTCATGGGTACCAGCACATAAAGGAATTGGAGGAAA TGAACAAATAGATAAATTAGTAAGTAAGGGAATCAGGAAAGTGCTGTTTCTAGATG **GAATAGAT**

17/23

TECH CENTER 1600/290

MAY 1.2 2003

SEQ ID NO:36

 $\begin{array}{c} GGCGGCATCGTGATCTACCAGTACATGGACGACCTGTACGTGGGCAGCGGCG\\ GC \end{array}$



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SEQ ID NO: 37

GGIVIYQYMDDLYVGSGG



19/23

12_5/1ZA (SEQ ID NO:45)

TGGAAGGGTTAATTTACTCCAGGAAAAGGCAAGAGATCCTTGATTTATGGGTCTATC ACACACAAGGCTACTTCCCTGATTGGCAAAACTACACACCGGGACCAGGGGTCAGA TATCCACTGACCTTTGGATGGTGCTTCAAGCTAGTGCCAGTTGACCCAAGGGAAGTA GAAGAGGCCAACGGAGGAGAAGACAACTGTTTGCTACACCCTATGAGCCAGTATGG AATGGATGATGAACACAAAGAAGTGTTACAGTGGAAGTTTGACAGCAGCCTAGCAC GCAGACACCTGGCCCGCGAGCTACATCCGGATTATTACAAAGACTGCTGACACAGA AGGGACTTTCCGCCTGGGACTTTCCACTGGGCGTTCCAGGGGGAGTGGTCTGGGCG GGACTGGGAGTGGCCAGCCCTCAGATGCTGCATATAAGCAGCGGCTTTTCGCCTGTA CTGGGTCTCTCTAGGTAGACCAGATCCGAGCCTGGGAGCTCTCTGTCTATCTGGGGA ACCCACTGCTTAGGCCTCAATAAAGCTTGCCTTGAGTGCTCTAAGTAGTGTGTGCCC ATCTGTTGTGACTCTGGTAACTCTGGTAACTAGAGATCCCTCAGACCCTTTGTGGT AGTGTGGAAAATCTCTAGCAGTGGCGCCCCGAACAGGGACTTGAAAGCGAAAGTGAG ACCAGAGAAGATCTCTCGACGCAGGACTCGGCTTGCTGAAGTGCACTCGGCAAGAG AGGAGAGAGATGGGTGCGAGAGCGTCAATATTAAGAGGGGGGAAAATTAGACAAAT GGGAAAAATTAGGTTACGGCCAGGGGGGAGAAAACACTATATGCTAAAACACCTA GTATGGGCAAGCAGAGAGCTGGAAAGATTTGCAGTTAACCCTGGCCTTTTAGAGAC ATCAGACGGATGTAGAC AAATAATAAAACAGCTACAACCAGCTCTTCAGA CAGGAACAGAGGAAATTAGATCATTATTTAACACAGTAGCAACTCTCTATTGTGTAC ATAAAGGGATAGATGTACGAGACACCAAGGAAGCCTTAGACAAGATAGAGGAGGA ACAAAACAAATGTCAGCAAAAAACACAGCAGGCGGAAGCGGCTGACAAAAAGGTC AGTCAAAATTATCCTATAGTGCAGAACCTCCAAGGCCAAATGGTACACCAGGCCAT ATCACCTAGAACCTTGAATGCATGGGTAAAAGTAATAGAGGAGAAGGCTTTTAGCC CAGAGGTAATACCCATGTTTACAGCATTATCAGAAGGAGCCACCCCACAAGATTTA AACACCATGTTAAATACAGTGGGGGGACATCAAGCAGCCATGCAAATGTTAAAAG ATACCATCAATGAGGAGGCTGCAGAATGGGATAGGTTACATCCAGTACATGCAGGG CCTGTTGCACCAGGCCAGATGAGAGAGCCAAGGGGAAGTGACATAGCAGGAACTA CTAGTACCCTTCAAGAACAAATAGCATGGATGACAAGTAACCCACCTATCCCAGTA CAGCCCTGTCAGCATTTTAGACATAAAACAAGGACCAAAGGAACCCTTTAGAGACT ATGTAGACCGGTTCTTCAAAACTTTAAGAGCTGAACAATCTACACAAGAGGTAAAA AATTGGATGACAGACACCTTGTTAGTCCAAAATGCGAACCCAGATTGTAAGACCATT TTAAGAGCATTAGGACCAGGGGCTTCATTAGAAGAAATGATGACAGCATGTCAGGG AGTGGGAGGACCTAGCCACAAAGCAAGAGTTTTGGCTGAGGCAATGAGCCAAGCAA ACAATACAAGTGTAATGATACAGAAAAGCAATTTTAAAGGCCCTAGAAGAGCTGTT AAATGTTTCAACTGTGGCAGGGAAGGGCACATAGCCAGGAATTGCAGGGCCCCTAG GAAAAGGGGCTGTTGGAAATGTGGAAAGGAAGGACACCAAATGAAAGACTGTACT GAGAGGCAGGCTAATTTTTTAGGGAAAATTTGGCCTTCCCACAAGGGGAGGCCAGG GAATTTCCTTCAGAGCAGACCAGAGCCAACAGCCCACCACTAGAACCAACAGCCC CACCAGCAGAGAGCTTCAAGTTCAAGGAGACTCCGAAGCAGGAGCCGAAAGACAG GGAACCTTTAACTTCCCTCAAATCACTCTTTGGCAGCGACCCCTTGTCTCAATAAAA

FIG. 16A



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GTAGCGGGCCAAACAAAGGAGGCTCTTTTAGATACAGGAGCAGATGATACAGTACT AGAAGAAATAAACTTGCCAGGAAAATGGAAACCAAAAATGATAGGAGGAATTGGA GGTTTTATCAAAGTAAGACAGTATGATCAAATACTTATAGAAATTTGTGGAAAAAGG GCTATAGGTACAGTATTAGTAGGACCTACACCTGTCAACATAATTGGAAGAAATCTG TTGACTCAGCTTGGATGCACACTAAATTTTCCAATTAGCCCCATTGAAACTGTACCA GTAAAATTAAAGCCAGGAATGGATGGCCCAAAGGTTAAACAATGGCCATTGACAGA ATTACAAAAATTGGGCCTGAAAATCCATATAACACTCCAGTATTTGCCATAAAGAAG AAGGACAGTACAAAGTGGAGAAAATTAGTAGATTTCAGGGAACTCAATAAAAGAAC TCAAGACTTTTGGGAAGTCCAATTAGGAATACCACACCCAGCAGGGTTAAAAAAGA AAAAATCAGTGACAGTACTGGATGTGGGAGATGCATATTTTTCAGTCCCTTTAGATG AGAGCTTCAGAAAATATACTGCATTCACCATACCTAGTATAAACAATGAAACACCA GGGATTAGATATCAATATATGTTCTTCCACAGGGATGGAAAGGATCACCAGCAA TATTCCAGAGTAGCATGACAAGAATCTTAGAGCCCTTTAGAACACAAAACCCAGAA GTAGTTATCTATCAATATATGGATGACTTATATGTAGGATCTGACTTAGAAATAGGG CAACATAGAGCAAAAATAGAGGAGTTAAGAGGACACCTATTGAAATGGGGATTTAC CACACCAGACAAGAACATCAGAAAGAACCCCCATTTCTTTGGATGGGGTATGAAC TCCATCCTGACAAATGGACAGTACAGCCTATACAGCTGCCAGAAAAGGAGAGCTGG ACTGTCAATGATATACAGAAGTTAGTGGGAAAGTTAAACTGGGCAAGTCAGATTTA CCCAGGGATTAAAGTAAGGCAACTGTGTAAACTCCTTAGGGGAGCCAAAGCACTAA CAGACATAGTGCCACTGACTGAAGAAGCAGAATTAGAATTGGCTGAGAACAGGGA AATTCTAAAAGAACCAGTACATGGAGTATATTATGACCCATCAAAAGATTTAATAG CTGAAATACAGAAACAGGGGAATGACCAATGGACATATCAAATTTACCAAGAACC ATTTAAAAATCTGAGAACAGGAAAGTATGCAAAAATGAGGACTGCCCACACTAATG ATGTGAAACAGTTAGCAGAGGCAGTGCAAAAGATAACCCAGGAAAGCATAGTAATA TGGGGAAAAACTCCTAAATTTAGACTACCCATCCCAAAAGAAACATGGGAGACATG GTGGTCAGACTATTGGCAAGCCACCTGGATTCCTGAGTGGGAGTTTGTCAATACCCC TCCCCTAGTAAAATTGTGGTACCAGCTGGAAAAAGAACCCATAGTAGGGGCAGAAA CTTTCTATGTAGATGGAGCAGCCAATAGGGAAACTAAAATAGGAAAAGCAGGGTAT GTCACTGACAAAGGAAGGCAGAAAGTTGTTTCCTTCACTGAAACAACAAATCAGAA GACTGAATTACAAGCAATTCAGCTAGCTTTGCAGGATTCAGGGCCAGAAGTAAACA TAGTAACAGACTCACAGTATGCATTAGGAATCATTCAAGCACAACCAGATAAGAGT GAATCAGAATTAGTCAGTCAAATAATAGAACAGTTGATAAAAAAGGAAAAAGTCTA CCTATCATGGGTACCAGCACATAAAGGAATTGGAGGAAATGAACAAGTAGACAAAT TAGTAAGTAGTGGAATCAGAAAAGTACTGTTTCTAGATGGAATAGATAAAGCTCAA GAAGAGCATGAAAAATATCACAGCAATTGGAGAGCAATGGCTAGTGAGTTTAATCT GCCACCCATAGTAGCAAAGGAAATAGTAGCCAGCTGTGATAAATGTCAGCTAAAAG GGGAAGCCATGCATGGACAAGTCGACTGTAGTCCAGGAATATGGCAATTAGACTGT ACACATTTAGAAGGAAAAATCATCCTAGTAGCAGTCCATGTAGCCAGTGGCTACAT GGAAGCAGAGGTTATCCCAGCAGAAACAGGACAAGAAACAGCATACTTATACTAA AATTAGCAGGAAGATGGCCAGTCAAAGTAATACATACAGATAATGGCAGTAATTTC ACCAGTACCGCAGTTAAGGCAGCCTGTTGGTGGGCAGATATCCAACGGGAATTTGG AATTCCCTACAATCCCCAAAGTCAAGGAGTAGTAGAATCCATGAATAAAGAATTAA

FIG. 16B



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AGAAAATCATAGGGCAAGTAAGAGATCAAGCTGAGCACCTTAAGACAGCAGTACAA ATGGCAGTATTCATTCACAATTTTAAAAGAAAAGGGGGGGATTGGGGGGTACAGTGC AGGGGAGAGAATAATAGACATAGCATCAGACATACAAACTAAAGAATTACAAA AACAAATTATAAAAATTCAAAATTTTCGGGTTTATTACAGAGACAGCAGAGACCCTA TTTGGAAAGGACCAGCCAAACTACTCTGGAAAGGTGAAGGGGCAGTAGTAATACAA GATAATAGTGATATAAAGGTAGTACCAAGAAGGAAAGCAAAAATCATTAAGGACTA TGGAAAACAGATGGCAGGTGCTGATTGTGTGGCAGGTAGACAGGATGAAGATTAGA CCCATTAGGAGATGCCAGGTTAGTAATAAAAACATATTGGGGTCTGCAGACAGGAG AAAGAGCTTGGCATTTGGGTCACGGAGTCTCCATAGAATGGAGATTGAGAAGATAT AGCACACAAGTAGACCCTGACCTGACAGACCAACTAATTCATATGCATTATTTTGAT TGTTTTGCAGAATCTGCCATAAGGAAAGCCATACTAGGACAGATAGTTAGCCCTAA GTGTGACTATCAAGCAGGACATAACAAGGTAGGATCTCTACAATACTTGGCACTGA CAGCATTGATAAAACCAAAAAAGATAAAGCCACCTCTGCCTAGTGTTAGGAAATTA GTAGAGGATAGATGGAACAAGCCCCAGAAGACCAGGGGCCGCAGAGGGAACCATA CAATGAATGGACACTAGAGCTTTTAGAAGAACTCAAGCAGGAAGCTGTCAGACACT TTCCTAGACCATGGCTCCATAACTTAGGACAACATATCTATGAAACCTATGGAGATA CTTGGACAGGAGTTGAAGCAATAATAAGAATCCTGCAACAATTACTGTTTATTCATT TCAGGATTGGGTGCCATCATAGCAGAATAGGCATTTTGCGACAGAGAAGAGCAAGA AATGGAGCCAATAGATCCTAACCTAGAACCCTGGAACCATCCAGGAAGTCAGCCTA AAACTGCTTGTAATGGGTGTTACTGTAAACGTTGCAGCTATCATTGTCTAGTTTGCTT TCAGAAAAAGGCTTAGGCATTTACTATGGCAGGAAGAAGCGGAGACAGCGACGAA AATAGTATATGTAATGTTAGATTTAACTGCAAGAATAGATTCTAGATTAGGAATAGG GAAAGAGCAGAAGACAGTGGCAATGAGAGCGAGGGGGATACTGAAGAATTATCGA CACTGGTGGATATGGGCATCTTAGGCTTTTGGATGCTAATGATGTGAATGTGAA GGGCTTGTGGGTCACAGTCTACTACGGGGTACCTGTGGGGAGAGAAGCAAAAACT GGCTACACATGCCTGTGTACCCACAGACCCCAACCCACAAGAAGTGATTTTGGGC AATGTAACAGAAAATTTTAACATGTGGAAAAATGACATGGTGGATCAGATGCAGG AAGATATAATCAGTTTATGGGATCAAAGCCTTAAGCCATGTGTAAAA1TGACCCCA CTCTGTGTCACTTTAAACTGTACAAATGCAACTGTTAACTACAATAATACCTCTAAA GACATGAAAAATTGCTCTTTCTATGTAACCACAGAATTAAGAGATAAGAAAAAGAA AGAAAATGCACTTTTTTATAGACTTGATATAGTACCACTTAATAATAGGAAGAATGG GAATATTAACAACTATAGATTAATAAATTGTAATACCTCAGCCATAACACAAGCCTG TCCAAAAGTCTCGTTTGACCCAATTCCTATACATTATTGTGCTCCAGCTGGTTATGCG CCTCTAAAATGTAATAAGAAATTCAATGGAATAGGACCATGCGATAATGTCAG CACAGTACAATGTACACATGGAATTAAGCCAGTGGTATCAACTCAATTACTGTTAAA TCAAAACAATAATAGTACATCTTAATGAATCTATAGAGATTAAATGTACAAGACC

FIG. 16C



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TGGCAATAATACAAGAAAGAGTGTGAGAATAGGACCAGGACAAGCATTCTATGCA ACAGGAGACATAATAGGAGATATAAGACAAGCACATTGTAACATTAGTAAAAATGA ATGGAATACAACTTTACAAAGGGTAAGTCAAAAATTACAAGAACTCTTCCCTAATA GTACAGGGATAAAATTTGCACCACACTCAGGAGGGGACCTAGAAATTACTACACAT AGCTTTAATTGTGGAGGAGAATTTTTCTATTGCAATACAACAGACCTGTTTAATAGT ACATACAGTAATGGTACATGCACTAATGGTACATGCATGTCTAATAATACAGAGCG CATCACACTCCAATGCAGAATAAAACAAATTATAAACATGTGGCAGGAGGTAGGAC GGACTACTATTAACACGTGATGGAGGAGATAATAATACTGAAACAGAGACATTCAG ACCTGGAGGAGGAGACATGAGGGACAATTGGAGAAGTGAATTATATAAATACAAG GTGGTAGAAATTAAACCATTAGGAGTAGCACCCACTGCTGCAAAAAGGAGAGTGGT GGAGAGAAAAAAGAGCAGTAGGAATAGGAGCTGTGTTCCTTGGGTTCTTGGGAG CAGCAGGAAGCACTATGGGCGCAGCATCAATAACGCTGACGGTACAGGCCAGACAA TTATTGTCTGGTATAGTGCAACAGCAAAGTAATTTGCTGAGGGCTATAGAGGCGCAA CAGCATATGTTGCAACTCACGGTCTGGGGCATTAAGCAGCTCCAGGCAAGAGTCCTG GCTATAGAGAGATACCTACAGGATCAACAGCTCCTAGGACTGTGGGGCTGCTCTGG AAAACTCATCTGCACCACTAATGTGCTTTGGAACTCTAGTTGGAGTAATAAAACTCA AAGTGATATTTGGGATAACATGACCTGGATGCAGTGGGATAGGGAAATTAGTAATT TGAAAAAGATTTACTAGCATTGGACAGGTGGAACAATCTGTGGAATTGGTTTAGCAT AACAAATTGGCTGTGGTATATAAAAATATTCATAATGATAGTAGGAGGCTTGATAG GTTTAAGAATAATTTTTGCTGTGCTCTCTCTAGTAAATAGAGTTAGGCAGGGATACT CACCCTTGTCATTGCAGACCCTTATCCCAAACCCGAGGGGGACCCGACAGGCTCGGA GGAATCGAAGAAGAAGGTGGAGAGCAAGACAGCAGCAGATCCATTCGATTAGTGA GCGGATTCTTGACACTTGCCTGGGACGACCTACGAAGCCTGTGCCTCTTCTGCTACC ACCGATTGAGAGACTTCATATTAATTGTAGTGAGAGCAGTGGAACTTCTGGGACAC AGTAGTCTCAGGGGACTGCAGAGGGGGGGGGGGAACCCTTAAGTATTTGGGGAGTCT TGTGCAATATTGGGGTCTAGAGTTAAAAAAAGAGTGCTATTAATCTGCTTGATACTAT AGCAATAGCAGTAGCTGAAGGAACAGATAGGATTCTAGAATTCATACAAAACCTTT GTAGAGGTATCCGCAACGTACCTAGAAGAATAAGACAGGGCTTCGAAGCAGCTTTG CAATAAAATGGGGGCAAGTGGTCAAAAAGCAGTATAATTGGATGGCCTGAAGTAA GAGAAAGAATCAGACGAACTAGGTCAGCAGCAGAGGGAGTAGGATCAGCGTCTCA AGACTTAGAGAAACATGGGGCACTTACAACCAGCAACACAGCCCACAACAATGCTG CTTGCGCCTGGCTGGAAGCGCAAGAGGAGGAAGGAGAAGTAGGCTTTCCAGTCAGA CCTCAGGTACCTTTAAGACCAATGACTTATAAAGCAGCAATAGATCTCAGCTTCTTT TTAAAAGAAAAGGGGGGACTGGAAGGGTTAATTTACTCCAAGAAAAGGCAAGAGAT CCTTGATTTGTGGGTTTATAACACACAAGGCTTCTTCCCTGATTGGCAAAACTACAC ACCGGGACCAGGGTCAGATTTCCACTGACCTTTGGATGGTACTTCAAGCTAGAGCC AGTCGATCCAAGGGAAGTAGAAGAGGCCAATGAAGGAGAAAACAACTGTTTACTAC ACCCTATGAGCCAGCATGGAATGGAGGATGAAGACAGAGAAGTATTAAGATGGAAG TTTGACAGTACGCTAGCACGCAGACACATGGCCCGCGAGCTACATCCGGAGTATTAC AAAGACTGCTGACACAGAAGGGACTTTCCGCTGGGACTTTCCACTGGGGCGTTCCAG GAGGTGTGGTCTGGGCGGGACAGGGGAGTGGTCAGCCCTGAGATGCTGCATATAAG CAGCTGCTTTTCGCCTGTACTGGGTCTCTCTAGGTAGACCAGATCTGAGCCCGGGAG

FIG. 16D



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CTCTCTGGCTATCTAGGGAACCCACTGCTTAAGCCTCAATAAAGCTTGCCTTGAGTG CCTTGAGTAGTGTGCCCGTCTGTTGTGTGACTCTGGTAACTAGAGATCCCTCAGA CCACTTGTGGTAGTGTGAAAATCTCTAGCA

FIG. 16E